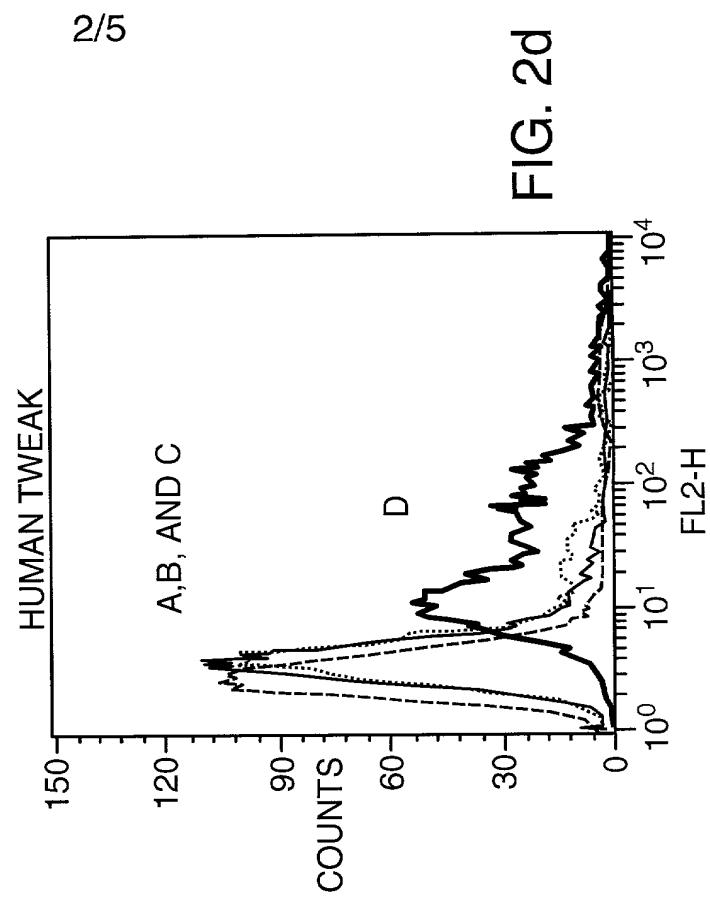
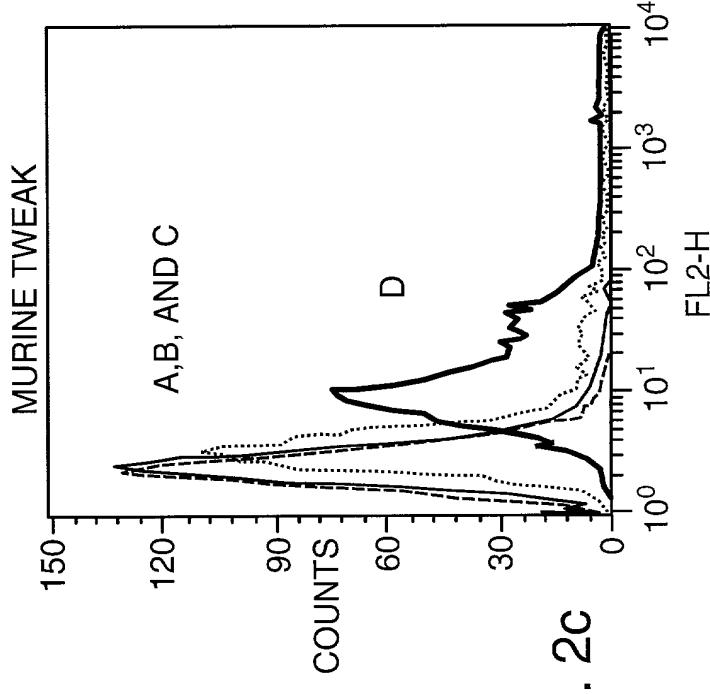
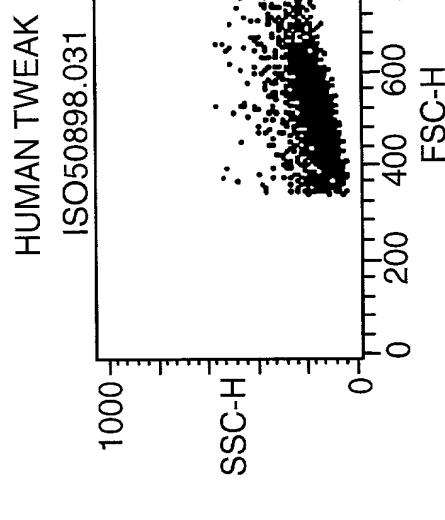
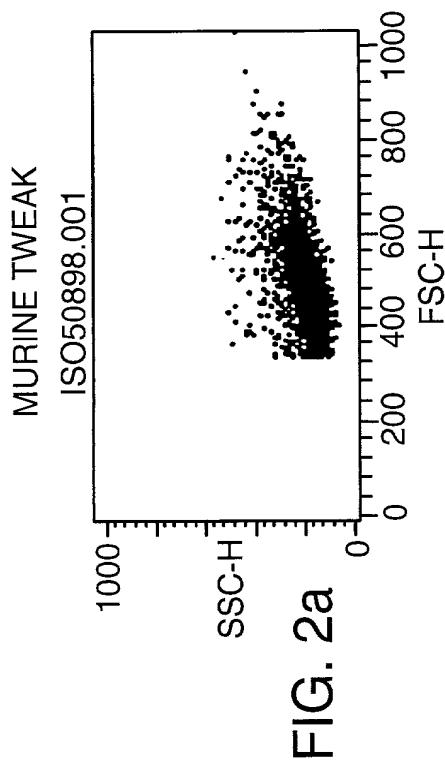


Human	1	MAARRSQRERRGRGEPTALLVPLAIGGLAAACIGLLAVSLSGRASL
Mouse	1	-----VLSGLAAACIGLLVVSLSGRASL
Human	51	SAQEPAAQEEELVAEEEDQDPSELNPQTEESQDPAPFLNRLVPRRSAPKGRK
Mouse	27	SAQEPSSQEEELTAEDRREPPELNQTEESQDVVPFLEQLVPRRSAPKGRK
Human	101	TRARRATAAHYEVHPRPGQDGQAQAGVTDGTVSGWEEEARINSSSPLRYNRQI
Mouse	77	APRARRATAAHYEVHPRPGQDGQAQAGVTDGTVSGWEEETKINSSSPLRYDRQI
Human	151	GEFIVTRAGLYLYCQVHFDEGKAVYLKLDLIVDGVLAIRCLEEFSATAA
Mouse	127	GEFTVTRAGLYLYCQVHFDEGKAVYLKLDLIVNGVLAIRCLEEFSATAA
Human	201	SSLGPGQLRLCQVSGLLALRPGSSLRIRTLPW AHLKAAPFLTYFGLFQVH
Mouse	177	SSSPGPQLRLCQVSGLLPLRPGSSLRIRTLPW AHLKAAPFLTYFGLFQVH

FIG. 1



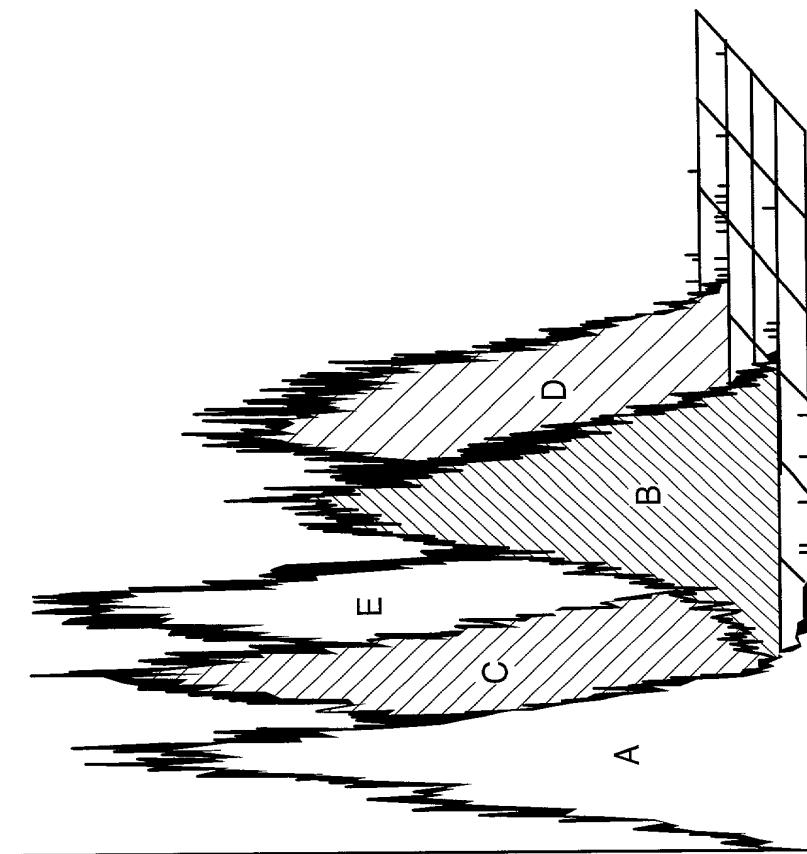


FIG. 3b

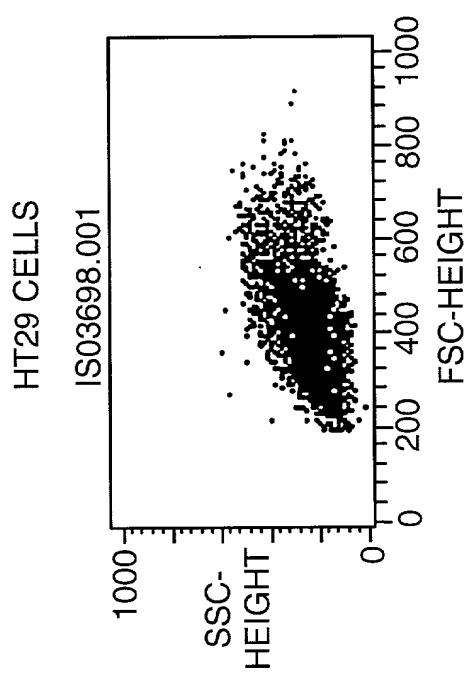


FIG. 3a

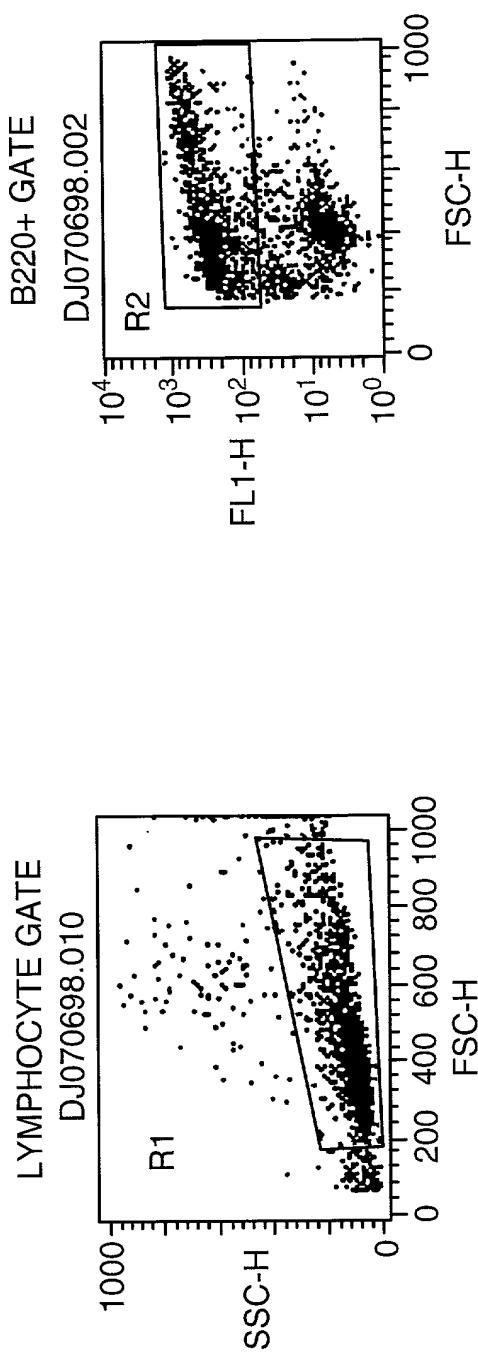
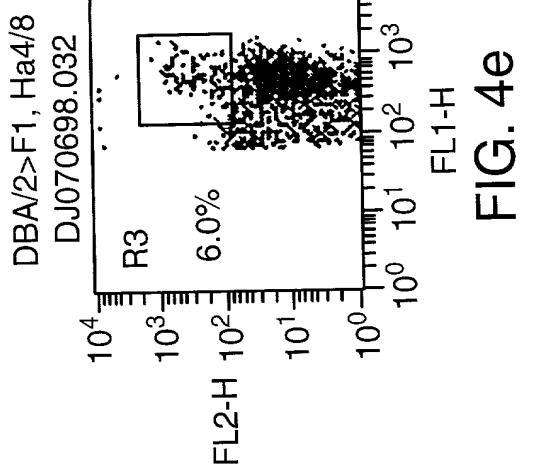
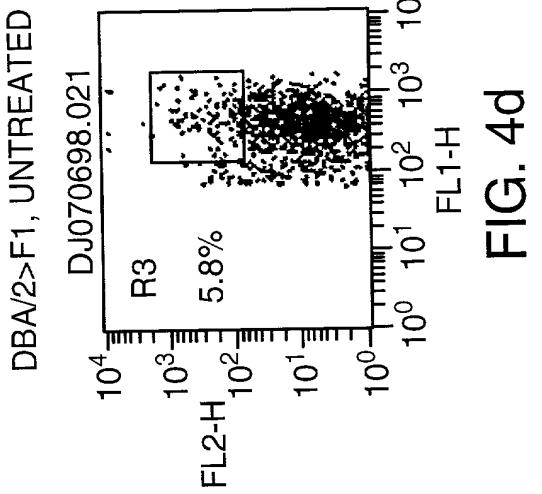
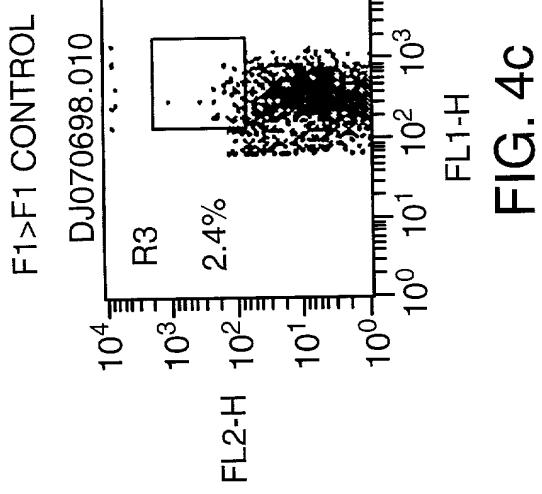


FIG. 4a

FIG. 4b



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